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RAW SEQUENCE LISTING

DATE: 05/31/2002

PATENT APPLICATION: PCT/US02/14570

TIME: 09:05:49

Input Set : A:\3198-WO.txt

Output Set: N:\CRF3\05312002\PU14570.raw

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	Glu .																9	0.1
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Input Set : A:\3198-WO.txt

66 gag aga ctc ggc ctg gag aga ctc ctgat ctg aga gag cag aga gag cag 508 67 Glu Arg Leu Gly Leu Glu Lys Leu Asp Leu Lys Arg Gln Lys Glu Gln 556 70 get ctg cgg gag gtg gag cac ctg aga gag tgc cag cag cag atg gct 556 71 Ala Leu Arg Glu Val Glu His Leu Lys Arg Cys Gln Gln Gln Met Ala 125 130 135 74 gag gac aag gcc tct gtg aaa gcc cag gtg ag tcc ttg ggg gag 604 160 115 125 74 gag gac aag gcc tct gtg aaa gcc cag gtg ag t ccc ttg ctc ggg gag 604 160 160 150 160	64		90					95					100					
67 Glu Arg Leu Gly Leu Glu Lys Leu Asp Leu Lys Arg Gln Lys Glu Gln 68 105 110 115 115 120 120 70 gct ctg cgg gag gtg gag cac ctg aag aga tgc cag cag cag atg gct 71 Ala Leu Arg Glu Val Glu His Leu Lys Arg Cys Gln Gln Gln Met Ala 72 125 130 135 74 gag gac aag gcc tct gtg aaa gcc cag gtg acg tcc ttg gct ggg gag 75 Glu Asp Lys Ala Ser Val Lys Ala Gln Val Thr Ser Leu Leu Gly Glu 76 140 145 150 78 ctg cag gag ag ag ag cag atg cyt gcg ac aca gag atg gcc gag 79 Leu Gln Glu Ser Gln Ser Arg Leu Glu Ala Ala Thr Lys Glu Cys Gln 80 155 82 gct ctg gag ggt cgg gcc cgg gcg gcc ag cag cag gcg cgg cag ctg 83 Ala Leu Glu Gly Arg Ala Arg Ala Ala Ser Glu Gln Ala Ala Thr Lys Glu Cys Gln 84 170 175 180 86 gag agt gag gag cgg cgc dc cag cag cag cag cag cgg cgg cag ctg 87 Glu Ser Glu Arg Glu Ala Leu Gln Gln His Ser Val Gln Val Asp 88 185 190 195 200 90 cag ctg cgc atg cag gag cag acg cag cag ccg cgc ctg cag cag cag ctg cag ctg cag ctg cag cag cag ctg cag ctg cag ctg cag cag cag ccg ctg cag cag cag cag cag cag cag ctg cag cag cag cag cag cag cag cag cag ca	66	gag	aga	ctc	ggc	ctg	gag	aag	ctc	gat	ctg	aag	agg	cag	aag	gag	cag	508
70																		
11 Ala Leu Arg Glu Val Glu His Leu Lys Arg Cys Gln Gln Gln Met Ala 135 125 130 135 130 135 135 136 135 136 1	68	105					110					115					120	
125	70	gct	ctg	cgg	gag	gtg	gag	cac	ctg	aag	aga	tgc	cag	cag	cag	atg	gct	556
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75 Glu Asp Lys Ala Ser Val Lys Ala Gln Val Thr Ser Leu Clu Glu Glu Clu Clu Glu Clu Glu Clu	72					125					130					135		
76 140 145 150 78 ctg cag gag agc cag agt cgc ttg gag gct gcc act ag gaa tgc cag 652 79 Leu Gln Glu Ser Gln Ser Arg Leu Glu Ala Ala Thr Lys Glu Cys Gln 652 80 155 160 165 82 gct ctg gag ggt cgg gcc cgg gcc agc gac agc gag cag cgc gcc agc gag cag c	74	gag	gac	aag	gcc	tct	gtg	aaa	gcc	cag	gtg	acg	tcc	ttg	ctc	ggg	gag	604
78 ctg cag gag agc cag agt cgc ttg gag gct gcc act aag gaa tgc cag 79 Leu Gln Glu Ser Gln Ser Arg Leu Glu Ala Ala Thr Lys Glu Cys Gln 80	75	Glu	Asp	Lys	Ala	Ser	Val	Lys	Ala	Gln	Val	Thr	Ser	Leu	Leu	Gly	Glu	
79 Leu Gln Glu Ser Gln Ser Arg Leu Glu Ala Ala Thr Lys Glu Cys Gln 80	76				140					145					150			
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83 Ala Leu Glu Gly Arg Ala Arg Ala Ala Ser Glu Gln Ala Arg Gln Leu 84 170 180 85 gag and gag cgc gag gcg ctg cag cag cag cac agc gtg cag gtg gac 748 87 Glu Ser Glu Arg Glu Ala Leu Gln Gln Gln His Ser Val Gln Val Asp 88 185 190 190 200 90 cag ctg cgc atg cag ggc cag agc gtg gag gcc gcg ctc cgc atg gag 796 91 Gln Leu Arg Met Gln Gly Gln Ser Val Glu Ala Leu Arg Met Glu 92 205 210 215 94 cgc cag gcc gcc tcg gag aga gag aga aga cgc gcg ctg cag ttg cag gtg gag 98 844 95 Arg Gln Ala Ala Ser Glu Glu Lys Arg Lys Leu Ala Gln Leu Gln Val 96 220 225 230 98 gcc tat cac cag ctc ttc caa gaa tac gac aac cac atc aag agc agc 99 Ala Tyr His Gln Leu Pre Gln Glu Tyr Asp Asn His Ile Lys Ser Ser 100 235 240 245 102 gtg gtg ggc agt gag cgg aag cga gag atg cag cag gag atg cag gag gag gag gag gag gag gag gag ga	80			155					160					165				
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87 Glu Ser Glu Arg Glu Ala Leu Gln Gln Gln His Ser Val Gln Val Asp 185 185 190 200 90 cag ctg cgc atg cag ggc cag agg ggg ggg gc gcg ctc cgc atg gag 91 Gln Leu Arg Met Gln Gly Gln Ser Val Glu Ala Ala Leu Arg Met Glu 92 205 210 215 94 cgc cag gcc gcc tcg gag gag aag agg agg ctg gcc cag ttg cag gtg 95 Arg Gln Ala Ala Ser Glu Glu Lys Arg Lys Leu Ala Gln Leu Gln Val 96 220 225 230 98 gcc tat cac cag ctc ttc caa gaa tac gac aac cac atc aag agc agc agc 99 Ala Tyr His Gln Leu Phe Gln Glu Tyr Asp Asn His Ile Lys Ser Ser 100 235 240 245 102 gtg gtg ggc agt gag cgg aag cga gg atg cgg gga atg ctg gac gat ctc aaa 940 103 Val Val Gly Ser Glu Arg Lys Arg Gly Met Gln Leu Gln Asp Leu Lys 104 250 255 260 106 cag cag cag ctc cag cag gcc gag gag gac ctg gga aaa cag gag gtg 107 Gln Gln Leu Gln Gln Ala Glu Glu Ala Leu Val Ala Lys Gln Glu Val 108 265 270 275 280 110 atc gat aag ctg aag gag gag gcc cag gag cac aac aag att gtg atg gag 111 Ile Asp Lys Leu Lys Glu Glu Ala Glu Gln His Lys Ile Val Met Glu 112 285 290 295 114 acc gtt ccg gtg ctg aag gcc cag gcg gat atc tac aag gcg gat ttc 1084 115 Thr Val Pro Val Leu Lys Ala Gln Ala Asp Ile Tyr Lys Ala Asp Phe 116 300 305 310 118 Cag gag ctg gag cag ctg gag cag ctg gag aag ctg ctg gac aac aag aac ag gag ctc 1132 129 ctg cag gad cag ctg cag gag cag ctg cag ag ag ag ctg ctg aag ag ag ag ctg ctg 118 Cag gag acg cag ctg aag gac aac cag aac aac aac ag gag gac ctc 1184 119 Gln Ala Glu Arg Gln Ala Arg Glu Lys Leu Ala Glu Lys Lys Glu Glu Leu 110 20 315 320 325 120 ctg cag gag cag ctg cag aag cag ctg cag aag ctg cag aac cac aac aac ac aac aac ac ac ac ac	84		170		-	_		175					180					
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92																		
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106 cag cag cag cag gag gag gcc ctg gtg gcc aaa cag gag gtg 988 107 Gln Gln Leu Gln Ala Glu Glu Ala Leu Val Leu Lys Glu Val 110 atc gat aag ctg aag gag gcc gag cac aag atc gtg gag 1036 111 11e Asp Lys Leu Lys Glu Glu Ala Glu His Lys Ile Val Met Glu 112 acc gtt ccg gtg ctg cag gcc gag cag gad atc tac atc gtg gag tac lus	103	Val	. Val	. Gly	Ser	Glu	Arg	Lys	Arg	Gly	Met	Gln	Leu	ı Glu	Asp	Leu	Lys	
107 Gln Gln Gln Leu Gln Gln Ala Glu Glu Ala Leu Val Ala Lys Gln Glu Val 108 265	104		250)				255					260)				
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110 atc gat aag ctg aag gag gag gcc gag cag cac aag att gtg atg gag 1036 111 Ile Asp Lys Leu Lys Glu Glu Ala Glu Gln His Lys Ile Val Met Glu 112	107	Gln	Glr	Leu	Gln	Gln	Ala	Glu	Glu	Ala	Leu	ı Val	. Ala	Lys	Gln	Glu	Val	
111 Ile Asp Lys Leu Lys Glu Glu Ala Glu Gln His Lys Ile Val Met Glu 295 114 acc gtt ccg gtg ctg aag gcc cag gcg gat atc tac aag gcg gac ttc 1084 115 Thr Val Pro Val Leu Lys Ala Gln Ala Asp Ile Tyr Lys Ala Asp Phe 310 118 cag gct gag agg cag gcc cgg gag aag ctg gcc gag aag ctc 1132 119 Gln Ala Glu Arg Gln Ala Arg Glu Lys Leu Ala Glu Lys Leu Ala Glu Lys Lys Glu Leu 20 120 ctg cag gag cag ctg gag cag ctg gag cag ctg gag agg ctg 325 122 ctg cag gag cag ctg gag cag ctg gag cag ctg cag agg gag tac agc aag ctg 325 123 Leu Gln Glu Gln Leu Glu Gln Leu Gln Arg Gln Arg Glu Tyr Ser Lys Leu Lys 124 124 330 c	108	265	ı				270)				275	i				280	
112	110	ato	gat	. aag	ctg	aag	gag	gag	gcc	gag	cag	cac	aag	att	. gtg	atg	gag	1036
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115 Thr Val Pro Val Leu Lys Ala Gln Ala Asp Ile Tyr Lys Ala Asp Phe 116	112	!				285					290	1				295		
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	Ser G	1v C1v	Dro	_	λla	λen	Gln	λen		Lou	C117	Clu	Clu		Dro	
179	.*	ту сту	FIO	ALG	Ата	тэр	GIII	чэћ	v a r	neu	Сту	GIU	GIU	267		
1,,			20					25					3.0		FIO	
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RAW SEQUENCE LISTING DATE: 05/31/2002 PATENT APPLICATION: PCT/US02/14570 TIME: 09:05:49

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	Tyr	ASP	ASII	HIS		гуѕ	ser	ser	val		GIY	ser	Glu	Arg	-	Arg		
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RAW SEQUENCE LISTING
PATENT APPLICATION: PCT/US02/14570

DATE: 05/31/2002 TIME: 09:05:49

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VERIFICATION SUMMARY

PATENT APPLICATION: PCT/US02/14570

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Input Set : A:\3198-WO.txt

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